

Title: NOVEL
MEGAKARYOCYTIC PROTEIN
TYROSINE KINASES
Inventor(s): Axel ULLRICH et al.
Appl. No.: 09/977,261

CTCGCTCCAAGTTGTCAGCCCGGACCGCTCGGGGTGTCAGCCCGCTCGCGAGCCCTCTCGGGCGGGCGGGCGGGCTCGGG 90

GGCCCCCTGAGCAGAAAACAGGAAGAACAGGCTCGGTCCAGTGGCAGCCAGCTCCCTACCTCCTGTGCCAGCCGCTGGCCTGTGGCA 180

GGCCATTCCCAGCGTCCCCGACTGTGACCACTTGCTCAGTGTGCCTCTCACCTGCCTCAGTTCCCTCTCGGGGGCGATGGCGGGCGAG 270
M A G R

GCTCTCGTTTCTCGCGGGCATTTACGGCTGTGATTCTGCTGAGGAAGTTCCCGGGTGAGCCCCGCTTCTCCGAGCCTGGCACC 360
G S L V S W R A F H G C D S A E E L P R V S P R F L R A W H

CCCCTCCGCTCTCAGCCAGGATGCCAAGAGCGCTGGGCCCCGGGACCCAGTGTATCACCAATCGGAGCACACCCGCCCAAGCCAG 450
P P P V S A R M P T R R W A P G T Q C I T K C E H T R P K P

GGGAGCTGCCCTTCGCAAGGGCGACGTGGTCACCATCTGGAGCCCTGGGAGAACAAGAGCTGGTACCGGCTCAAGCACCACACCACTG 540 SH 3
G E L A F R K G D V V T I L E A C E N K S W Y R V K H H T S

GACAGGAGCGCTGCTGGCAGCTGGGGCGCTGGGGAGCGGAGGCCCTCTCCGAGACCCCAAGCTCAGCCTCATGCCGTGGTTCCAGC 630
G Q E G L L A A G A L R E R E A L S A D P K L S L M P W F H

GGAAGATCTCGGGCCAGGAGCTGTCCAGCAGCTGCAGCCCTCCCGAGGATGGGTGTTCTGGTGGGGAGTCCGCGGCCACCCGGCG 720
G K I S G Q E A V Q Q L Q P P E D G L F L V R E S A R H P G SH 2

ACTACGTCCTGTGGTGAGCTTTGCCCGGACGTATCCACTACCGGTGCTGCACCGCGACGGCCACCTCACAATCGATAGGCCGTGT 810
D Y V L C V S F G R D V I H Y R V L H R D G H L T I D E A V

TCTTCTGCAACCTCATGACATGGTGGAGCATTACAGCAAGGACAAGCGGCTATCTGCACCAAGCTGGTGAGACCAAGCGGAAACAGC 900
F F C N L M D M V E H Y S K D K G A I C T K L V R P K R K H

FIG.1A

Title: NOVEL
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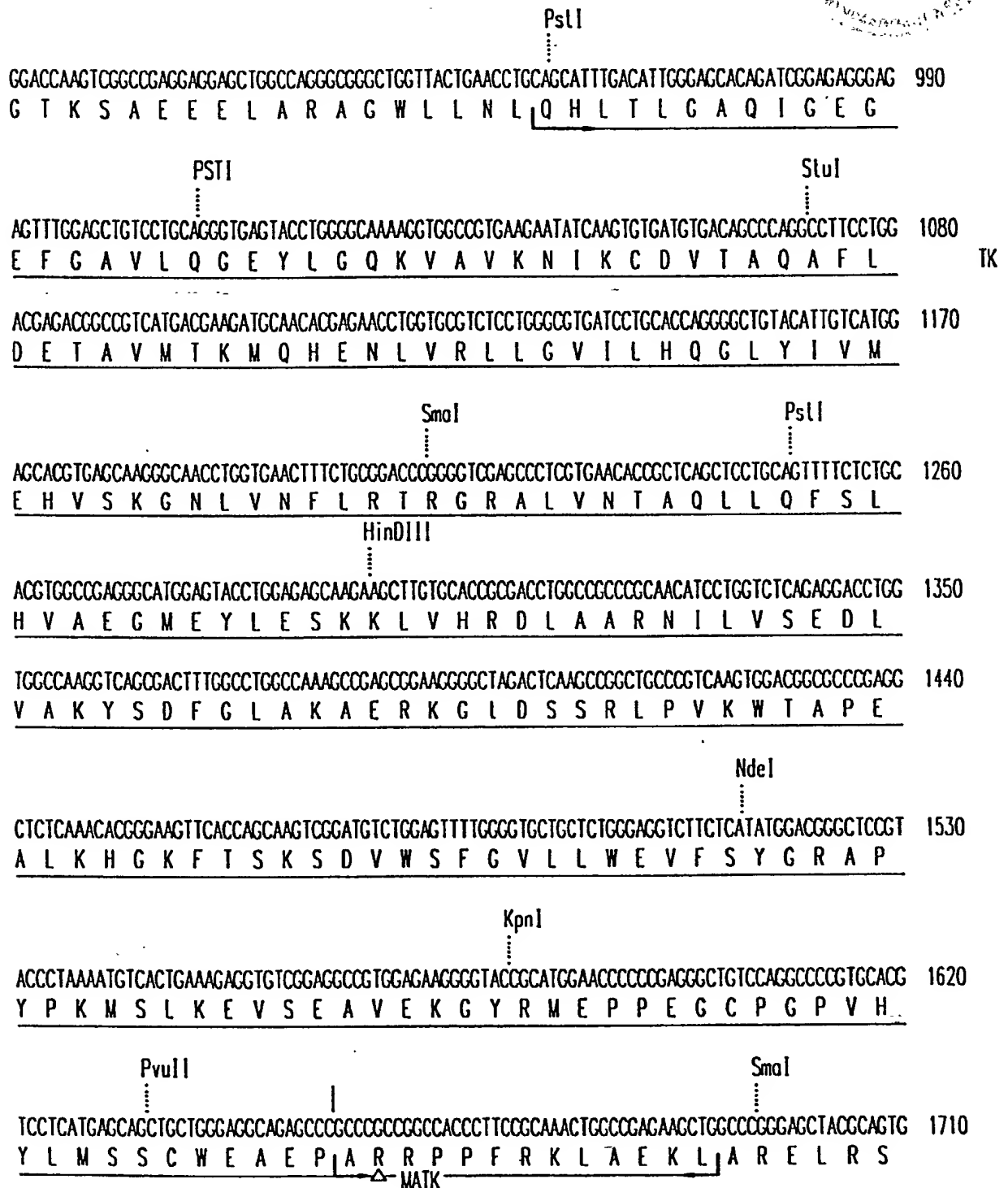
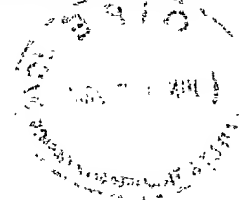


FIG.1B

Title: NOVEL
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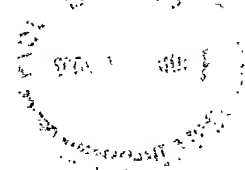
977261-060202



CAGGTGCCCCAGCCTCCGTCTCAGGGCAGGAGCCGACGGCTCCACCTGCCCCGAAGCCAGGAGCCCTGACCCACCCGGTGGGGCCCT 1800
A G A P A S V S G Q D A D G S T S P R S Q E P
TGGCCCCAGAGACCGAGAGAGTGAGAGTGCGCGGTGGGGGCACTGACCAGGCCCAAGGAGGTCCAGGCGGCAAGTCATCCTCCTGG 1890
TGCCACAGCAGGGGCTGGCCACGTAGGGGCTCTGCGCGCCCGTGGACACCCACACCTGCCAAGGATGATGCCCCGATAAAGACCG 1980
ATTCTAAGGACTCTAAAAA 2000

FIG.1C

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CCGCTTTTTCCTTAGAGCTTGAGAGTCAAAG AGGACCCACATGTATACTTCGGCTCTAGCGAGT AGGATGATAATATGGATACA 90
M D T

AAATCTATTCTAGAAGAACTTCTTCTCAAAAGATCACAGCAAAAGAAGAAAATGTCACCAAAATAATTACAAGAACGGCTTTTGTGTTG 180
K S I L E E L L L K R S Q Q K K K M S P N N Y K E R L F V L

ACCAAAACAAACCTTTCCTACTATGAATATGACAAATGAAAAGGGCAGCAGAAAAGGATCCATTGAAATTAAGAAAATCAGATGTGTG 270
T K T N L S Y Y E Y D K M K R G S R K G S I E I K K I R C V

GAGAAAGTAAATCTCGAGGAGCAGACGCCCTGTAGAGAGACAGTACCCATTTAGATTGTCTATAAGATGGGCTTCTCTATGTCTATGCA 360
E K V N L E E Q T P V E R Q Y P F Q I V Y K D G L L Y V Y A PH

TCAATGAAGAGAGCCGAAGTCAGTGGTTGAAAGCATTACAAAAGAGATAAGGGTAACCCCCACCTGCTGCTCAAGTACCATAGTGGG 450
S N E E S R S Q W L K A L Q K E I R G N P H L L V K Y H S G

TTCTTGTGCGACCGGAAGTTCCTGTGTGCCAGCAGAGCTGTAAGCAGCCCCAGGATGTACCTCTCGGAAGCATATGCTAATCTGCAT 540
F F V D G K F L C C Q Q S C K A A P G C T L W E A Y A N L H

ACTGCAGTCAATGAAGAGAAACACAGAGTTCACCTTCCCAGACAGAGTGTGAAGATACCTCGGCAGTTCTGTCTCAAAATGGAT 630
T A V N E E K H R V P T F P D R V L K I P R A V P V L K M D

GCACCATCTTCAAGTACCACTCTAGCCCAATATGACAACGAATCAAAGAAAACTATGGCTCCCAGCCACCATCTTCAAGTACCACTCTA 720
A P S S S T T L A Q Y D N E S K K N Y G S Q P P S S S T S L SH3

GCGCAATATGACAGCAACTCAAAGAAAATCTATGGCTCCCAGCCAACTTCAACATGCAGTATATTCCAAGGAAGACTTCCCTGACTGG 810
A Q Y D S N S K K I Y G S Q P N F N M Q Y I P R E D F P D W

TGCAAGTAAGAAAAGTGAAGTAGCAGCAGCAGTGAAGATGTTGCAAGCAGTAACCAAAAAGAAAGAAATGTGAATCACACCACCTCA 900
W Q V R K L K S S S S S E D V A S S N Q K E R N V N H T T S

AAGATTTTCATCGGAATTCCTGAGTCAAGTTCATCTGAAGAGAGCAAAACCTCGATGATTATGACTGCTTTCCTGCTAACATCTCCAGA 990
K I S W E F P E S S S S E E E E N L D D Y D W F A G N I S R

TCACAATCTGAACAGTTACTCAGACAAAAGCGAAAAGAGGAGCATTATGCTTAGAAATTCAGCCAAGTGGGAATGTACACAGTGTCC 1080
S Q S E Q L L R Q K G K E G A F M V R N S S Q V G M Y T V S SH4

TTATTTAGTAAGCTGTGAATGATAAAAAAGGAAGTGTCAACATTACCAGTGCATACAAATGCTGAGAACAAATTATACCTGGCAGAA 1170
L F S K A V N D K K G T V K H Y H V H T N A E N K L Y L A E

FIG.2A

7261-000702

AACTACTGTTTGGATTCCATTCCAAAGCTTATTCATTATCATCAACACAATTACGAGGCATGATCACACGGCTCCGCCACCCCTGTGTCA 1260
N Y C F D S I P K L I H Y H Q H N S A G M I T R L R H P V S

ACAAAGGCCAACAAGCTCCCCACTCTGTGTCCCTCGGAAATGGAATCTCGGAAGTAAAAGAGAAGAGATTACCTTGTTCAGGAGCTG 1350
T K A N K V P D S V S L G N G I W E L K R E E I T L L K E L

GGAAGTGGCCAGTTTGGAGTGGTCCAGCTGGGCAAGTGAAGGGCAGTATGATGTTGCTGTTAAGATGATCAAGGAGGCTCCATGTCA 1440
G S G Q F G V V Q L G K W K G Q Y D V A V K M I K E G S M S

GAAGATGAATTCCTTCAGGAGGCCAGACTATGATGAAACTCAGCCATCCCAAGCTGGTTAAATTCTATCGAGTGTGTTCAAAGCAATAC 1530
E D E F F Q E A Q T M M K L S H P K L V K F Y G V C S K E Y

CCCATATACATAGTGAATATATAAGCAATGGCTGCTTGTGAATTACCTGAGGAGTACCGAAAAGGACTTGAACCTTCCAGCTC 1620 TK
P I Y I V T E Y I S N G C L L N Y L R S H G K G L E P S Q L

TTAGAAATGTCTACGATGTCTGTGAAGGCATGGCCTTCTTGGAGAGTCACCAATTCATACACGGGAGTTGGCTGCTCGTAAGTGTG 1710
L E M C Y D V C E G M A F L E S H Q F I H R D L A A R N C L

GTGGACAGAGATCTCTGTGTGAAGTATCTGACTTTGGAATGACAAGGTATGTTCTTGATGACCAGTATGTCAGTTCAGTCCGAACAAAG 1800
V D R D L C V K V S D F G M T R Y V L D D Q Y V S S V G T K

TTTCCAGTCAAGTGGTCAAGTCCAGAGGTGTTTCACTTCAATACAGCAGCAAGTCAAGCTATGGGCATTTGGGATCCTGATGTGG 1890
F P V K W S A P E V F H Y F K Y S S K S D V W A F G I L M W

GAGGTGTTCAAGCTGGGGAAGCAGCCCTATGACTTGTATGACAACCTCCAGGTGGTTCTGAAGGTCTCCAGGGCCACAGCTTTACCGG 1980
E V F S L G K Q P Y D L Y D N S Q V V L K V S Q G H R L Y R

CCCCACCTGGCATCGGACACCATCTACCAGATCATGTACAGCTGCTGGCAGGAGCTTCCAGAAAAGCGTCCACATTTACGCAACTCCTG 2070
P H L A S D T I Y Q I M Y S C W H E L P E K R P T F Q Q L L

TCTTCCATTGAACCACTTCGGGAAAAAGACAAGCATTGAAGAAGAAATTAGGAGTGTGATAAGAATGAATATAGATGCTGCCAGCATT 2160
S S I E P L R E K D K H .

TTCATTCAITTTAAGGAAAGTAGCAAGGCATAATGTAATTTAGCTAGTTTTTAATAGTGTCTCTGTATTGTCTATTATTTAGAAATGAA 2250

CAAGGCAGGAAACAAAAGATTCCCTTGAAATTTAGGTCAAATTAGTAATTTTGTATTGCTGCCCTGATATAACACTTTCCAGCCTATA 2340

GCAGAAGCACATTTTCAGACTGCAATATAGAGACTGTGTTATGTGTAAGACTGAGCAGAACTGAAAAATTACTTATTGGATATTCATT 2430

CTTTTCTTTATATTGTCATTGTGACACAATTAATATACTACCAAGTACAAAAAAAAAAAAAAAAAAAAA 2500

FIG.2B

CCGGACTGGTCGAAAGACAGGAACAGACTTGAACAGGGGAGAGCTCTGGCGAAACGAAGACGTGGAGGTTTTACCAGGGATAAGAAG 90
AAAGACACCTTCTAGTGAGCAGCTGCCAGCTCTGCTCAGTTTTGCCTCGGGTAGCACCTCCAGCCACAGAAAGCAAGCCGTAAG 180
TCTCTCCAGGTAGGACTTGCTGCAACCCAGCTGCTGGACTGATCTGAAACGGGACTTTGCATACTCTCCGAAGTATGGTGAGTTGGTCT 270
M V S W C
GACTTCAAAGTTGCTCGTGAAGGAAGATAAGGTGGATCGCAGAGACTAAGGGGAGAGGAGAAGCCCTGCTCTTCTCCCCACCAAG 360
GCACAATGAGCAACATCTGTCAGAGCTCTCGGAGTACCTAGAACCCTATCTCCCTGTTTGTCACGGAGGCAGACAAGTCAACCGTGA 450
M S N I C Q R L W E Y L E P Y L P C L S T E A D K S T V
TTGAAATCCAGGGCGCTTTGCTCTCCCAGTCACAGAGGCATGCCACTACTTTGTCGCTTTGTTTGATTACCAGGCTCGGACTGCTG 540
I E N P G A L C S P Q S Q R H G H Y F V A L F D Y Q A R T A
AGGACTTGAGCTTCGAGCAGGTGACAACTTCAAGTTCTGGACACTTTGCATGAGGGCTGGTGGTTTCCAGACACTTGGAGAAAAGAC 630 SH 3
E D L S F R A G D K L Q V L D T L H E G W W F A R H L E K R
GAGATGGCTCCAGTCAGCAACTACAAGCTATATTCCTTCTAAGTACGTGGCTGAGGACAGAAGCCTACAGGCAGAGCCGTTCTTTG 720
R D G S S Q Q L Q G Y I P S N Y V A E D R S L Q A E P W F F
GAGCAATCCGAAGATCAGATGCAGAGAAACAATATTATATTCAGAAAACAAGACCGGTTCTTTCTAATCAGAGAAAGTGAAGCCAAA 810
G A I G R S D A E K Q L L Y S E N K T G S F L I R E S E S Q SH 2
AAGGAGAATTCTCTCTTTAGATGGAGCAGTTGTAACACTACAGAATTAAGACTGGATGAAGGGGATTTTTTCTCAGGC 900
K G E F S L S V L D G A V V K H Y R I K R L D E G G F F L T
GAAGAAGAATCTTTCAACACTGAACGAATTTGTGAGCCACTACACCAAGACAAGTGACGGCCTGTGTGTCAGCTGGGAAACCATGCT 990
R R R I F S T L N E F V S H Y T K T S D G L C V K L G K P C
TAAAGATCCAGGTCCCAGCTCCATTTGATTTGCTGATAAAACCGTGGACCAATGGGAGATAGACCGCAACTCCATACAGCTTCTGAAGC 1080
L K I Q V P A P F D L S Y K T V D Q W E I D R N S I Q L L K
GATTGGGATCTGGTCAGTTTGGCGAAGTATGGGAAGGCTGTGGAACAATACCACTCCAGTAGCAGTGAAACATTAAACAGGTTCAA 1170
R L G S G Q F G E V W E G L W N N T T P V A V K T L K P G S
TGGATCCAAATGACTTCTGAGGGAGGCACAGATAATGAAGAACCTAAGACATCCAAAGCTTATCCAGCTTTATGCTGTTTGCACCTTAG 1260
M D P N D F L R E A Q I M K N L R H P K L I Q L Y A V C T L

FIG.3



		<u>MKK1</u>	<u>MKK2</u>
HUMAN			
MEG/ERYTH	MEG-01	+++	+++
	K562	++	+
	MO7E	++	+
	HEL	+++	++
MYELO/MAC	KG-1	+	++
	HL-60	+	+
	TF-1	+	+
B-CELL	ALL-1	-	+
	RAJI	-	-
	DAUDI	-	-
T-CELL	MOLT-3	-	-
	JURKAT	-	-
EPITHELIAL	HELA	-	-
RODENT			
	BM	+	+++
	SPLEEN	+++	+
	THYMUS	-	-
	LIVER	-	-
	BRAIN	+	-
RAT NEURAL	P19	+	-

FIG.4

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IN 977261 .060702

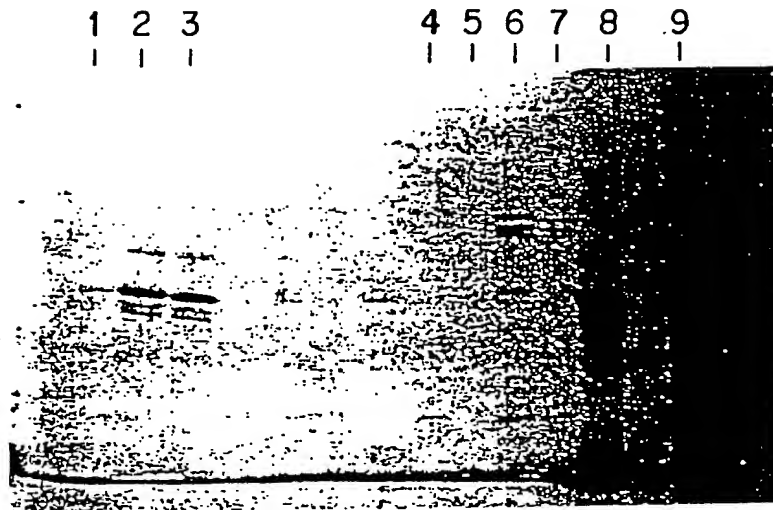
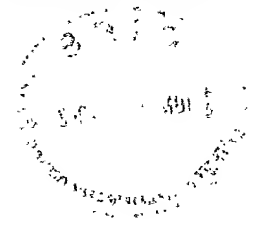


FIG. 5

Title: NOVEL
MEGAKARYOCYTIC PROTEIN
TYROSINE KINASES
Inventor(s): Axel ULLRICH et al.
Appl. No.: 09/977,261

09/977,261 .060702

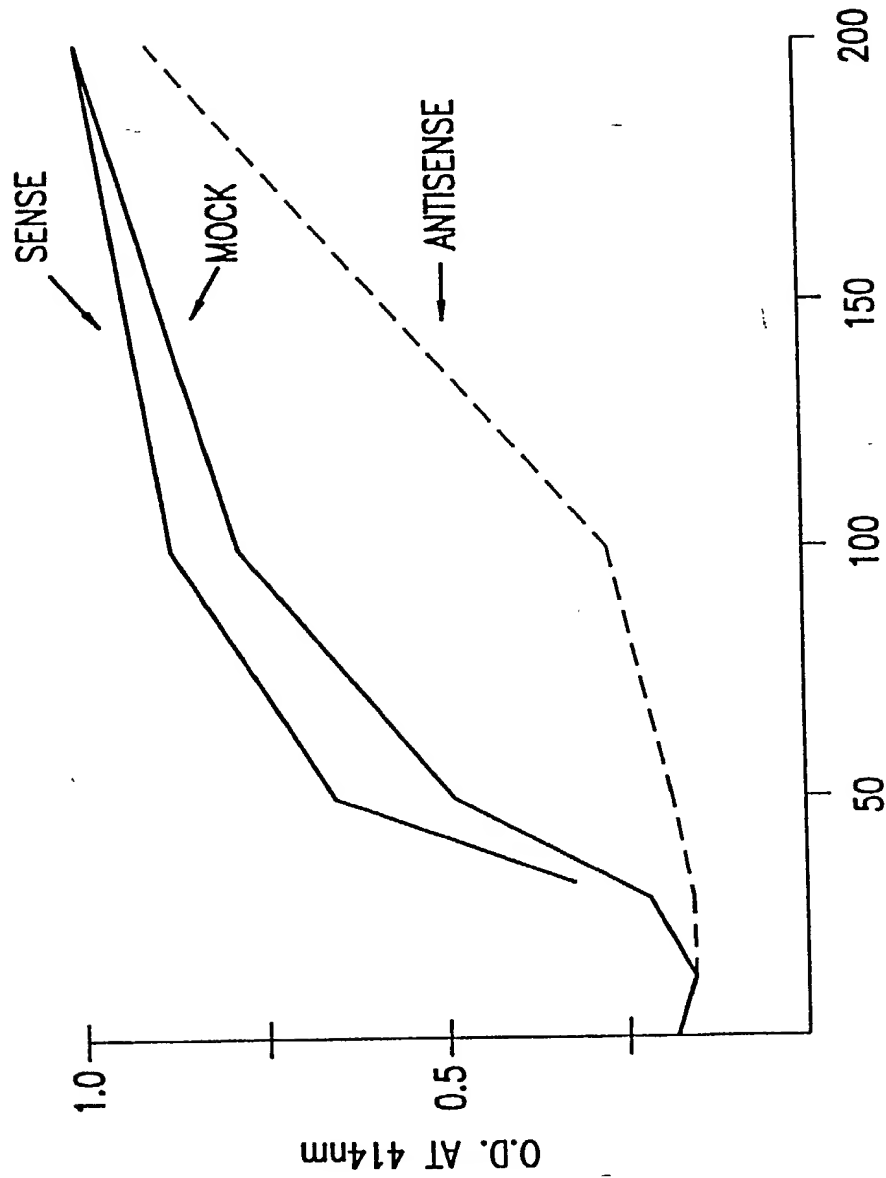


FIG. 6A

Title: NOVEL
MEGAKARYOCYTIC PROTEIN
TYROSINE KINASES
Inventor(s): Axel ULLRICH et al.
Appl. No.: 09/977,261

977261.060702



MKKI PROTEIN EXPRESSION

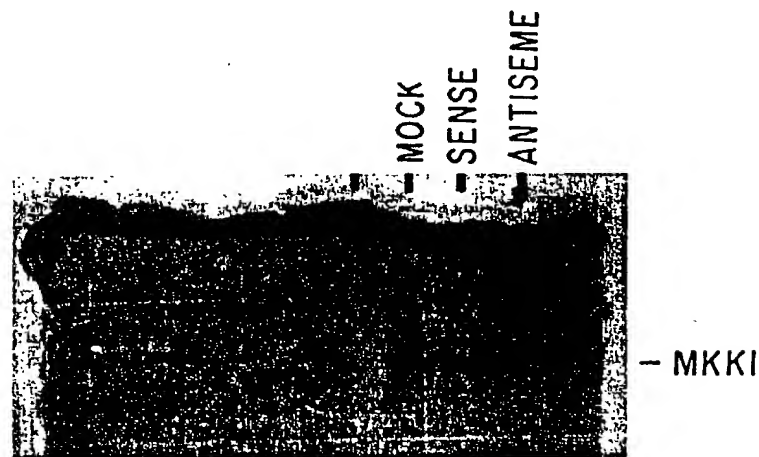


FIG. 6B

Title: NOVEL
MEGAKARYOCYTIC PROTEIN 977261.060702
TYROSINE KINASES
Inventor(s): Axel ULLRICH et al.
Appl. No.: 09/977,261

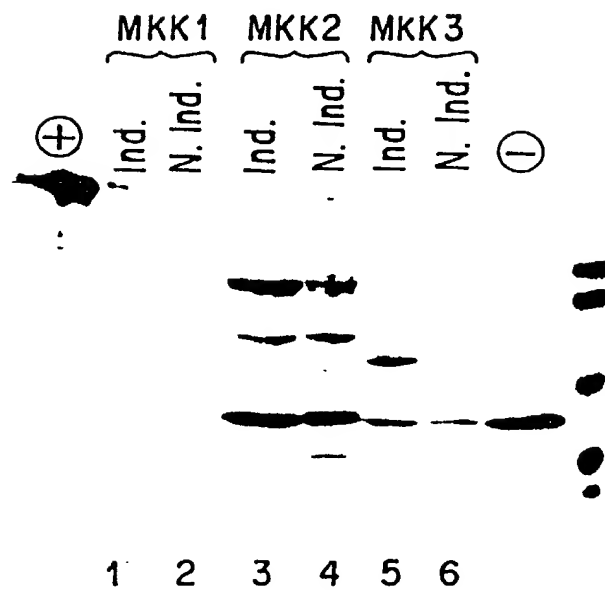
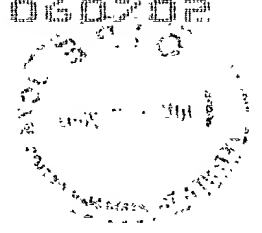


FIG. 7

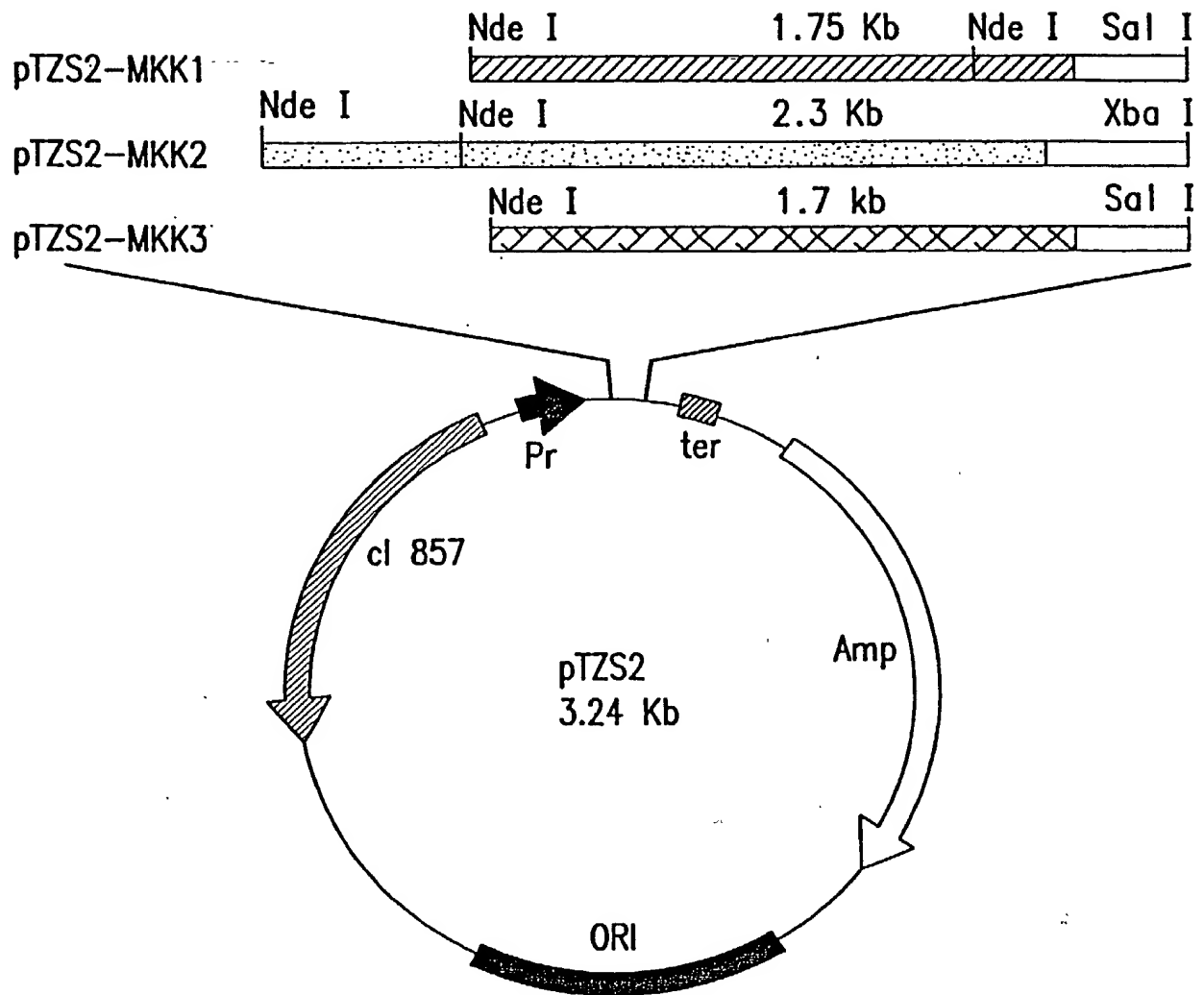


FIG.8

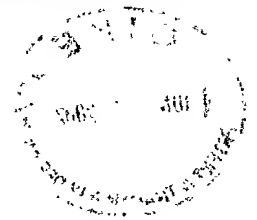
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 MEGAKARYOCYTIC PROTEIN 9977261.060702
 TYROSINE KINASES
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1	M A G R G S L V S W R A F H G C D S A E E L P R V S P R F L	MKK1 aa
1	M S A I Q A A - - - - -	hCSK (JH0559)
31	R A W H P P P V S A R M P T R R W A P G T O C I T K C E H T	MKK1 aa
8	- - - - - W P S G T E C I A K Y N F H	hCSK (JH0559)
61	R P K P G E L A F R K G D V V T I L E A C E N K S W Y R V K	MKK1 aa
22	G T A E Q D L P F C K G D V L T I V A V T K D P N W Y K A K	hCSK (JH0559)
91	H H T S G Q E G L L A A G A L R E R E A L S A D P K L S L M	MKK1 aa
52	N K V - G R E G I I P A N Y V Q K R E G V K A G T K L S L M	hCSK (JH0559)
121	P W F H G K I S G Q E A V Q Q L Q P P E D G L F L V R E S A	MKK1 aa
81	P W F H G K I T R E Q A E R L L Y P P E T G L F L V R E S T	hCSK (JH0559)
151	R H P G D Y V L C V S F G R D V I H Y R V L H R D G H L T I	MKK1 aa
111	N Y P G D Y T L C V S C D G K V E H Y R I M Y H A S K L S I	hCSK (JH0559)
181	D E A V F F C N L M D M V E H Y S K D K G A I C T K L V R P	MKK1 aa
141	D E E V Y F E N L M Q L V E H Y T S D A D G L C T R L I K P	hCSK (JH0559)
211	K R K H G T K S A E E E L A R A G W L L N L Q H L T L G A Q	MKK1 aa
171	K V M E G T V A A Q D E F Y R S G W A L N M K E L K L L Q T	hCSK (JH0559)
241	I G E G E F G A V L Q G E Y L G Q K V A V K N I K C D V T A	MKK1 aa
201	I G K G E F G D V M L G D Y R G N K V A V K C I K N D A T A	hCSK (JH0559)
271	Q A F L D E T A V M T K M Q H E N L V R L L G V I L H Q - -	MKK1 aa
231	Q A F L A E A S V M T Q L R H S N L V Q L L G V I V E E K G	hCSK (JH0559)
299	G L Y I V M E H V S K G N L V N F L R T R G R A L V N T A Q	MKK1 aa
261	G L Y I V T E Y M A K G S L V D Y L R S R G R S V L G G D C	hCSK (JH0559)
329	L L Q F S L H V A E G M E Y L E S K K L V H R D L A A R N I	MKK1 aa
291	L L K F S L D V C E A M E Y L E G N N F V H R D L A A R N V	hCSK (JH0559)
359	L V S E D L V A K V S D F G L A K A E R K G L D S S R L P V	MKK1 aa
321	L V S E D N V A K V S D F G L T K E A S S T Q D T G K L P V	hCSK (JH0559)

FIG.9A

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389	K W T A P E A L K H G K F T S K S D V W S F G V L L W E V F	MKK1 aa
351	K W T A P E A L R E K K F S T K S D V W S F G I L L W E I Y	hCSK (JH0559)
419	S Y G R A P Y P K M S L K E V S E A V E K G Y R M E P P E G	MKK1 aa
381	S F G R V P Y P R I P L K D V V P R V E K G Y K M D A P D G	hCSK (JH0559)
449	C P G P V H V L M S S C W E A E P A R R P P F R K L A E K L	MKK1 aa
411	C P P A V Y E V M K N C W H L D A A M R P S F L Q L R E Q L	hCSK (JH0559)
479	A R E L R S A G A P A S V S G Q D A D G S T S P R S Q E P	MKK1 aa
441	E H - - - - - I K T H E L H - - - - - L	hCSK (JH0559)

FIG.9B

1	M	D	T	K	S	I	L	E	E	L	L	K	R	S	Q	Q	K	K	K	M	S	P	N	N	Y	K	E	R	L	MKK2	aa		
1	M	A	A	-	V	I	L	E	S	I	F	L	K	R	S	Q	Q	K	K	K	T	S	P	L	N	F	K	K	R	L	hAtk	(X58957)	
1	M	N	N	F	I	L	L	E	E	Q	L	I	K	K	S	Q	Q	K	R	R	T	S	P	S	N	F	K	V	R	F	hTKT	(L10717)	
1	M	M	V	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	mTec	(X5663)	
31	F	V	L	T	K	T	N	L	S	Y	Y	E	-	-	Y	D	K	M	K	R	G	S	R	K	G	S	I	E	I	K	MKK2	aa	
30	F	L	L	T	V	H	K	L	S	Y	Y	E	Y	D	F	E	R	G	R	G	S	K	K	G	S	I	D	V	E	hAtk	(X58957)		
31	F	V	L	T	K	A	S	L	A	Y	F	E	D	R	-	-	H	G	K	K	R	T	L	K	G	S	I	E	L	S	hTKT	(L10717)	
4	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	mTec	(X5663)	
59	K	I	R	C	V	E	K	V	N	L	E	E	Q	T	P	V	E	R	Q	-	-	-	-	-	-	-	-	-	-	-	-	MKK2	aa
60	K	I	T	C	V	E	T	V	V	P	E	K	N	P	P	P	E	R	Q	I	P	R	R	G	E	E	S	S	E	M	hAtk	(X58957)	
59	R	I	K	C	V	E	I	V	K	S	D	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	hTKT	(L10717)	
4	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	mTec	(X5663)	
78	-	-	-	-	-	-	-	-	-	-	-	Y	P	F	Q	I	V	Y	K	D	G	L	L	Y	V	Y	A	S	N	E	E	MKK2	aa
90	E	Q	I	S	I	I	E	R	F	P	Y	P	F	Q	V	V	Y	D	E	G	P	L	Y	V	F	S	P	T	E	E	hAtk	(X58957)	
70	-	-	I	S	I	P	C	H	Y	K	Y	P	F	Q	V	V	H	D	N	Y	L	L	Y	V	F	A	P	D	R	E	hTKT	(L10717)	
4	-	-	-	-	-	-	-	-	-	-	S	F	P	V	K	I	N	F	H	S	S	P	-	-	-	-	-	-	-	-	Q	mTec	(X5663)
98	S	R	S	Q	W	L	K	A	L	Q	K	E	I	R	G	N	P	H	L	L	V	K	Y	H	S	G	F	F	V	D	MKK2	aa	
120	L	R	K	R	W	I	H	Q	L	K	N	V	I	R	Y	N	S	D	L	V	Q	K	Y	H	P	C	F	W	I	D	hAtk	(X58957)	
98	S	R	Q	R	W	V	L	A	L	K	E	E	T	R	N	N	S	L	V	P	K	Y	H	P	N	F	W	M	D	hTKT	(L10717)		
17	S	R	D	R	W	V	K	K	L	K	E	E	T	K	N	N	N	I	M	I	K	Y	H	P	K	F	W	A	D	mTec	(X5663)		
128	G	K	F	L	C	C	Q	Q	S	C	K	A	A	P	G	C	T	L	W	E	A	Y	A	N	L	H	T	A	V	N	MKK2	aa	
150	G	Q	Y	L	C	C	S	Q	T	A	K	N	A	M	G	C	Q	I	L	E	N	R	N	G	S	L	K	P	G	S	hAtk	(X58957)	
128	G	K	W	R	C	C	S	Q	L	E	K	L	A	T	G	C	A	Q	Y	D	-	-	-	-	-	-	-	-	-	P	hTKT	(L10717)	
47	G	S	Y	Q	C	C	R	Q	T	E	K	L	A	P	G	C	E	K	Y	N	L	F	E	S	S	I	-	-	-	-	mTec	(X5663)	
158	E	E	K	H	R	V	P	T	F	P	D	R	V	L	K	I	P	R	A	V	P	V	L	K	M	D	A	P	S	S	MKK2	aa	
180	S	H	R	K	T	K	K	P	L	P	P	-	-	-	-	T	P	E	E	D	Q	I	L	K	K	P	L	P	P	E	hAtk	(X58957)	
149	T	K	N	A	S	K	K	P	L	P	P	-	-	-	-	T	P	E	D	N	R	-	-	-	-	-	-	-	-	hTKT	(L10717)		
73	-	-	-	-	-	R	K	T	L	P	P	-	-	-	-	A	P	E	-	-	-	-	I	K	K	R	R	P	P	-	mTec	(X5663)	
188	S	T	T	L	A	Q	Y	D	N	E	S	K	K	N	Y	G	S	Q	P	P	S	S	S	T	S	L	A	Q	Y	D	MKK2	aa	
206	P	A	A	A	P	V	S	T	S	E	L	K	K	-	-	-	-	-	-	-	-	-	-	V	V	A	L	Y	D	hAtk	(X58957)		
166	-	-	-	R	P	L	W	E	P	E	E	T	V	-	-	-	-	-	-	-	-	-	-	V	I	A	L	Y	D	hTKT	(L10717)		
89	P	P	I	P	P	E	E	E	N	T	E	E	I	-	-	-	-	-	-	-	-	-	-	-	V	V	A	M	Y	D	mTEC	(X5663)	

FIG.10A

Title: NOVEL
MEGAKARYOCYTIC PROTEIN
TYROSINE KINASES
Inventor(s): Axel ULLRICH et al.
Appl. No.: 09/977,261

7/26/1 060702

218	S N S K K I Y G S Q P N F N M Q Y I P R E D F P - D W W Q V	MKK2 aa
225	Y M P M N A N D L Q L R K G D E Y F I L E E S N L P W W R A	hAtk (X58957)
182	Y Q T N D P Q E L A L R R N E E Y C L L D S S E I H W W R V	hTKT (L10717)
108	F Q A T E A H D L R L E R G Q E Y I I L E K N D L H W W R A	mTec (X5663)
247	R K L K S S S S S E D V A S S N Q K E R N V N H T T S K I S	MKK2 aa
255	R D - - K N G Q E G Y I P S N Y V T E - A - - - - -	hAtk (X58957)
212	Q D - - R N G H E G Y V P S S Y L V E K S - - - - -	hTKT (L10717)
138	R D - - K -	mTec (X5663)
277	W E F P E S S S S E E E E N L D D Y D W F A G N I S R S Q S	MKK2 aa
273	- - - - - E D S I E M Y E W Y S K H M T R S Q A	hAtk (X58957)
231	- - - - - P N N L E T Y E W Y N K S I S R D K A	hTKT (L10717)
141	- - - - - Y G W Y C R N T N R S K A	mTec (X5663)
307	E Q L L R Q K G K E G A F M V R N S S Q V G M Y T V S L F S	MKK2 aa
292	E Q L L K Q E G K E G G F I V R D S S K A G K Y T V S V F A	hAtk (X58957)
250	E K L L L D T G K E G A F M V R D S R T A G T Y T V S V F T	hTKT (L10717)
154	E Q L L R T E D K E G G F M V R D S S Q P G L Y T V S L Y T	mTec (X5663)
337	K - A V N D K K G T V K H Y H V H - - T N A E N K L Y L A E	MKK2 aa
322	K S T - G D P Q G V T R H Y V V - - C S T P Q S Q Y Y L A E	hAtk (X58957)
280	K A V V S E N N P C I K H Y H I K E T N D N P K R Y Y V A E	hTKT (L10717)
184	K F G - G E G S S G F R H Y H I K E T A T S P K K Y Y L A E	mTec (X5663)
364	N Y C F D S I P K L I H Y H Q H N S A G M I T R L R H P V S	MKK3 aa
349	K H L F S T I P E L I N Y H Q H N S A G L I S R L K Y P V S	hAtk (X58957)
310	K Y V F D S I P L L I N Y H Q H N G G G L V T R L R Y P V C	hTKT (L10717)
213	K H A F G S I P E I I E Y H K H N A A G L V T R L R Y P V S	mTec (X5663)
394	T K A N K V P D S V S L G N G I W E L K R E E I T L L K E L	MKK2 aa
379	Q Q N K N A P S T A G L G Y G S W E I D P K D L T F L K E L	hAtk (X58957)
340	F G R Q K A P V T A G L R Y G K W V I D P S E L T F V Q E I	hTKT (L10717)
243	T K G K N A P T T A G F S Y D K W E I N P S E L T F M R E L	mTec (X5663)
424	G S G Q F G V V Q L G K W K G Q Y D V A V K M I K E G S M S	MKK2 aa
409	G T G Q F G V V K Y G K W R G Q Y D V A I K M I K E G S M S	hAtk (X58957)
370	G S G Q F G L V H L G Y W L N K D K V A I K T I R E G A M S	hTKT (L10717)
273	G S G L F G V V R L G K W R A Q Y K V A I K A I R E G A M C	mTec (X5663)

FIG.10B

454	E D E F F Q E A Q T M M K L S H P K L V K F Y G V C S K E Y	MKK2 aa
439	E D E F I E E A K V M M N L S H E K L V Q L Y G V C T K Q R	hA tk (X58957)
400	E E D F I E E A E V M M K L S H P K L V Q L Y G V C L E Q A	hTKT (L10717)
303	E E D F I E E A K V M M K L T H P K L V Q L Y G V C T Q Q K	mTec (X5663)
484	P I Y I V T E Y I S N G C L L N Y L R S H G K G L E P S Q L	MKK2 aa
469	P I F I I T E Y M A N G C L L N Y L R E M R H R F Q T Q Q L	hA tk (X58957)
430	P I C L V F E F M E H G C L S D Y L R T Q R G L F A A E T L	hTKT (L10717)
333	P I Y I V T E F M E R G C L L N F L R Q R Q G H F S R D M L	mTec (X5663)
514	L E M C Y D V C E G M A F L E S H Q F I H R D L A A R N C L	MKK2 aa
499	L E M C K D V C E A M E Y L E S K Q F L H R D L A A R N C L	hA tk (X58957)
460	L G M C L D V C E G M A Y L E E A C V I H R D L A A R N C L	hTKT (L10717)
363	L S M C Q D V C E G M E Y L E R N S F I H R D L A A R N C L	mTec (X5663)
544	V D R D L C V K V S D F G M T R Y V L D D Q Y V S S V G T K	MKK2 aa
529	V N D Q G V V K V S D F G L S R Y V L D D E Y T S S V G S K	hA tk (X58957)
490	V G E N Q V I K V S D F G M T R F V L D D Q Y T S S T G T K	hTKT (L10717)
393	V N E A G V V K V S D F G M A R Y V L D D Q Y T S S S G A K	mTec (X5663)
574	F P V K W S A P E V F H Y F K Y S S K S D V W A F G I L M W	MKK2 aa
559	F P V R W S P P E V L M Y S K F S S K S D I W A F G V L M W	hA tk (X58957)
520	F P V K W A S P E V F S F S R Y S S K S D V W S F G V L M W	hTKT (L10717)
423	F P V K W C P P E V F N Y S R F S S K S D V W S F G V L M W	mTec (X5663)
604	E V F S L G K Q P Y D L Y D N S Q V V L K V S Q G H R L Y R	MKK2 aa
589	E I Y S L G K M P Y E R F T N S E T A E H I A Q G L R L Y R	hA tk (X58957)
550	E V F S E G K I P Y E N R S N S E V V E D I S T G F R L Y K	hTKT (L10717)
453	E T F T E G R M P F E K N T N Y E V V T M V T R G H R L H R	mTec (X5663)
634	P H L A S D T I Y Q I M Y S C W H E L P E K R P T F Q Q L L	MKK2 aa
619	P H L A S E K V Y T I M Y S C W H E K A D E R P T F K I L L	hA tk (X58957)
580	P R L A S T H V Y Q I M N H C W K E R P E D R P A F S R L L	hTKT (L10717)
483	P K L A T K Y L Y E V M L R C W Q E R P E G R P S F E D L L	mTec (X5663)
664	S S I E P L R E K D K H	MKK2 aa
649	S N I L D V M D E E S	hA tk (X58957)
610	R Q L A E I A E S - - - - G L	hTKT (L10717)
513	R T I D E L V E C E E T F G R	mTec (X5663)

FIG.10C

Inventor(s): Axel ULLRICH et al.
Appl. No.: 09/977,261

MKK3 MPI aa
 hF yn
 eYrk

MKK3 MP1 oo
hFyn
cYrk
hSrc
hYes
hFgr
hLyn
hHck
hLck
mB1k

MKK3 MP1 aa
hFyn
cYrk
hSrc
hYes
hFgr
hLyn
hHck
hLck
mB1k

MKK3 MP1 aa
hFyn
cYrk
hSrc
hYes
hFgr
hLyn
hHck
hLck
mB1k

FIG. 11A

Title: NOVEL
MEGAKARYOCYTIC PROTEIN
TYROSINE KINASES

Inventor(s): Axel ULLRICH et al.

Appl. No.: 09/977,261

69	L	Q	V	L	D	T	L	H	E	G	W	W	F	A	R	H	L	E	K	R	R	D	G	S	S	Q	Q	L	Q	G	MKK3 MPI aa
109	F	Q	I	L	N	S	S	E	G	D	W	W	E	A	R	S	L	T	T	G	E	T	G	-	-	-	-	-	-	-	hFyn
108	F	H	I	I	N	N	T	E	G	D	W	W	E	A	R	S	L	S	S	G	A	T	G	-	-	-	-	-	-	-	cYrk
111	L	Q	I	V	N	N	T	E	G	D	W	W	L	A	H	S	L	S	T	G	Q	T	G	-	-	-	-	-	-	-	hSrc
118	F	Q	I	I	N	N	T	E	G	D	W	W	E	A	R	S	I	A	T	G	K	N	G	-	-	-	-	-	-	-	hYes
104	F	H	I	L	N	N	T	E	G	D	W	W	E	A	R	S	L	S	S	G	K	T	G	-	-	-	-	-	-	-	hFgr
90	M	K	V	L	E	E	H	-	G	E	W	W	K	A	K	S	L	L	T	K	K	E	G	-	-	-	-	-	-	-	hLyn
84	M	V	V	L	E	E	S	-	G	E	W	W	K	A	R	S	L	A	T	R	K	E	G	-	-	-	-	-	-	-	hHck
88	L	R	I	L	E	Q	S	-	G	E	W	W	K	A	Q	S	L	T	T	G	Q	E	G	-	-	-	-	-	-	-	hLck
79	L	Q	V	L	R	S	T	-	G	D	W	W	L	A	R	S	L	V	T	G	R	E	G	-	-	-	-	-	-	-	mBlk
99	Y	I	P	S	N	Y	V	A	E	D	R	S	L	Q	A	E	P	W	F	F	G	A	I	G	R	S	D	A	E	K	MKK3 MPI aa
132	Y	I	P	S	N	Y	V	A	P	V	D	S	I	Q	A	E	E	W	Y	F	G	K	L	G	R	K	D	A	E	R	hFyn
131	Y	I	P	S	N	Y	V	A	P	V	D	S	I	Q	A	E	E	W	Y	F	G	K	I	G	R	K	D	A	E	R	cYrk
134	Y	I	P	S	N	Y	V	A	P	S	D	S	I	Q	A	E	E	W	Y	F	G	K	I	T	R	R	E	S	E	R	hSrc
141	Y	I	P	S	N	Y	V	A	P	A	D	S	I	Q	A	E	E	W	Y	F	G	K	M	G	R	K	D	A	E	R	hYes
127	C	I	P	S	N	Y	V	A	P	V	D	S	I	Q	A	E	E	W	Y	F	G	K	I	G	R	K	D	A	E	R	hFgr
112	F	I	P	S	N	Y	V	A	K	L	N	T	L	E	T	E	E	W	F	F	K	D	I	T	R	K	D	A	E	R	hLyn
106	Y	I	P	S	N	Y	V	A	R	V	D	S	L	E	T	E	E	W	F	F	K	G	I	S	R	K	D	A	E	R	hHck
110	F	I	P	F	N	F	V	A	K	A	N	S	L	E	P	E	P	W	F	F	K	N	L	S	R	K	D	A	E	R	hLck
101	Y	V	P	S	N	F	V	A	P	V	E	T	L	E	V	E	K	W	F	F	R	T	I	S	R	K	D	A	E	R	mBlk
129	Q	L	L	Y	S	E	N	K	T	G	S	F	L	I	R	E	S	E	S	Q	K	G	E	F	S	L	S	V	L	D	MKK3 MPI aa
162	Q	L	L	S	F	G	N	P	R	G	T	F	L	I	R	E	S	E	T	T	K	G	A	Y	S	L	S	I	R	D	hFyn
161	Q	L	L	C	H	G	N	C	R	G	T	F	L	I	R	E	S	E	T	T	K	G	A	Y	S	L	S	I	R	D	cYrk
164	L	L	L	N	A	E	N	P	R	G	T	F	L	V	R	E	S	E	T	T	K	G	A	Y	C	L	S	V	S	D	hSrc
171	L	L	L	N	P	G	N	Q	R	G	I	F	L	V	R	E	S	E	T	T	K	G	A	Y	S	L	S	I	R	D	hYes
157	Q	L	L	S	P	G	N	P	Q	G	A	F	L	I	R	E	S	E	T	T	K	G	A	Y	S	L	S	I	R	D	hLyn
142	Q	L	L	A	P	G	N	S	A	G	A	F	L	I	R	E	S	E	T	L	K	G	S	F	S	L	S	V	R	D	hHck
136	Q	L	L	A	P	G	N	M	L	G	S	F	M	I	R	D	S	E	T	T	K	G	S	Y	S	L	S	V	R	D	hHck
140	Q	L	L	A	P	G	N	T	H	G	S	F	L	I	R	E	S	E	S	T	A	G	S	F	S	L	S	V	R	D	hLck
131	Q	L	L	A	P	M	N	K	A	G	S	F	L	I	R	E	S	E	S	N	K	G	A	F	S	L	S	V	K	D	mBlk
159	-	-	-	-	-	G	A	V	V	K	H	Y	R	I	K	R	L	D	E	G	G	F	F	L	T	R	R	R	I	F	MKK3 MPI aa
192	W	D	D	M	K	G	D	H	V	K	H	Y	K	I	R	K	L	D	N	G	G	Y	Y	I	T	T	R	A	Q	F	hFyn
191	W	D	E	A	K	G	D	H	V	K	H	Y	K	I	R	K	L	D	S	G	G	Y	Y	I	T	T	R	A	Q	F	cYrk
194	F	D	N	A	K	G	L	N	V	K	H	Y	K	I	R	K	L	D	S	G	G	F	Y	I	T	S	R	T	Q	F	hSrc
201	W	D	E	I	R	G	D	N	V	K	H	Y	K	I	R	K	L	D	N	G	G	Y	Y	I	T	T	R	A	Q	F	hYes
187	W	D	Q	T	R	G	D	H	V	K	H	Y	K	I	R	K	L	D	M	G	G	Y	Y	I	T	T	R	V	Q	F	hFgr
172	F	D	P	V	H	G	D	V	I	K	H	Y	K	I	R	S	L	D	N	G	G	Y	Y	I	S	P	R	I	T	F	hLyn
166	Y	D	P	R	Q	G	D	T	V	K	H	Y	K	I	R	T	L	D	N	G	G	F	Y	I	S	P	R	S	T	F	hHck
170	F	D	Q	N	Q	G	E	V	V	K	H	Y	K	I	R	N	L	D	N	G	G	F	Y	I	S	P	R	I	T	F	hLck
161	I	T	T	-	Q	G	E	V	V	K	H	Y	K	I	R	S	L	D	N	G	G	Y	Y	I	S	P	R	I	T	F	mBlk

FIG.11B

Title: NOVEL
MEGAKARYOCYTIC PROTEIN
TYROSINE KINASES
Inventor(s): Axel ULLRICH et al.
Appl. No.: 09/977,261

09977261 .060702

184	S	T	L	N	E	F	V	S	H	Y	T	K	T	S	D	G	L	C	V	K	L	G	K	P	C	L	K	I	Q	V	MKK3 MPI	oo
222	E	T	L	Q	Q	L	V	Q	H	Y	S	E	R	A	A	G	L	C	C	R	L	V	V	P	C	H	K	G	M	-	hFyn	
221	D	T	I	Q	Q	L	V	Q	H	Y	I	E	R	A	A	G	L	C	C	R	L	A	V	P	C	P	K	G	T	-	cYrk	
224	N	S	L	Q	Q	L	V	A	Y	Y	S	K	H	A	D	G	L	C	H	R	L	T	T	V	C	P	T	S	K	-	hSrc	
231	D	T	L	Q	K	L	V	K	H	Y	T	E	H	A	D	G	L	C	H	K	L	T	T	V	C	P	T	V	K	-	hYes	
217	N	S	V	Q	E	L	V	Q	H	Y	M	E	V	N	D	G	L	C	N	L	L	I	A	P	C	T	I	M	K	-	hFgr	
202	P	C	I	S	D	M	I	K	H	Y	Q	K	Q	A	D	G	L	C	R	R	L	E	K	A	C	I	S	P	K	-	hLyn	
196	S	T	L	Q	E	L	V	D	H	Y	K	K	G	N	D	G	L	C	Q	K	L	S	V	P	C	M	S	S	K	-	hHck	
200	P	G	L	H	E	L	V	R	H	Y	T	N	A	S	D	G	L	C	T	R	L	S	R	P	C	Q	T	Q	K	-	hLck	
190	P	T	L	Q	A	L	V	Q	H	Y	S	K	K	G	D	G	L	C	Q	K	L	T	L	P	C	V	N	L	A	-	mBik	

214	P	A	P	F	D	L	S	Y	K	T	V	D	Q	W	E	I	D	R	N	S	I	Q	L	L	K	R	L	G	S	G	MKK3 MPI	oo
251	P	R	L	T	D	L	S	V	K	T	K	D	V	W	E	I	P	R	E	S	L	Q	L	I	K	R	L	G	N	G	hFyn	
250	P	K	L	A	D	L	S	V	K	T	K	D	V	W	E	I	P	R	E	S	L	Q	L	L	Q	K	L	G	N	G	cYrk	
253	P	Q	T	Q	G	L	A	-	-	-	K	D	A	W	E	I	P	R	E	S	L	R	L	E	V	K	L	G	Q	G	hSrc	
260	P	Q	T	Q	G	L	A	-	-	-	K	D	A	W	E	I	P	R	E	S	L	R	L	E	V	K	L	G	Q	G	hYes	
246	P	Q	T	L	G	L	A	-	-	-	K	D	A	W	E	I	S	R	S	S	I	T	L	E	R	R	L	G	T	G	hFgr	
231	P	Q	-	-	-	-	K	P	W	D	K	D	A	W	E	I	P	R	E	S	I	K	L	V	K	R	L	G	A	G	hLyn	
225	P	Q	-	-	-	-	K	P	W	E	K	D	A	W	E	I	P	R	E	S	L	K	L	E	K	K	L	G	A	G	hHck	
229	P	Q	-	-	-	-	K	P	W	W	E	D	E	W	E	V	P	R	E	T	L	K	L	V	E	R	L	G	A	G	hLck	
219	P	K	-	-	-	-	N	L	W	A	Q	D	E	W	E	I	P	R	Q	S	L	K	L	V	R	K	L	G	S	G	mBik	

244	Q	F	G	E	V	W	E	G	L	W	N	N	T	T	P	V	A	V	K	T	L	K	P	G	S	M	D	P	N	D	MKK3 MPI	oo
281	Q	F	G	E	V	W	M	G	T	W	N	G	N	T	K	V	A	I	K	T	L	K	P	G	T	M	S	P	E	S	hFyn	
280	Q	F	G	E	V	W	M	G	T	W	N	G	T	T	K	V	A	V	K	T	L	K	P	G	T	M	S	P	E	A	cYrk	
280	C	F	G	E	V	W	M	G	T	W	N	G	T	T	R	V	A	I	K	T	L	K	P	G	T	M	S	P	E	A	hSrc	
287	C	F	G	E	V	W	M	G	T	W	N	G	T	T	K	V	A	I	K	T	L	K	P	G	T	M	M	P	E	A	hYES	
273	C	F	G	D	V	W	L	G	T	W	N	G	S	T	K	V	A	V	K	T	L	K	P	G	T	M	S	P	K	A	hFgr	
257	Q	F	G	E	V	W	M	G	Y	Y	N	N	S	T	K	V	A	V	K	T	L	K	P	G	T	M	S	V	Q	A	hLyn	
251	Q	F	G	E	V	W	M	A	T	Y	N	K	H	T	K	V	A	V	K	T	M	K	P	G	S	M	S	V	E	A	hHck	
255	Q	F	G	E	V	W	M	G	Y	Y	N	G	H	T	K	V	A	V	K	S	L	K	Q	G	S	M	S	P	D	A	hLck	
245	Q	F	G	E	V	W	M	G	Y	Y	K	N	N	M	K	V	A	T	K	T	L	K	E	G	T	M	S	P	E	A	mBik	

274	F	L	R	E	A	Q	I	M	K	N	L	R	H	P	K	L	I	Q	L	Y	A	V	C	T	L	E	D	P	I	Y	MKK3 MPI	oo
311	F	L	E	E	A	Q	I	M	K	K	L	K	H	D	K	L	V	Q	L	Y	A	V	V	S	-	E	E	P	I	Y	hFyn	
310	F	L	E	E	A	Q	I	M	K	R	L	R	H	D	K	L	V	Q	L	Y	A	V	V	S	-	E	E	P	I	Y	cYrk	
310	F	L	Q	E	A	Q	V	M	K	K	L	R	H	E	K	L	V	Q	L	Y	A	V	V	S	-	E	E	P	I	Y	hSrc	
317	F	L	Q	E	A	Q	I	M	K	K	L	R	H	D	K	L	V	P	L	Y	A	V	V	S	-	E	E	P	I	Y	hYes	
303	F	L	E	E	A	Q	V	M	K	L	L	R	H	D	K	L	V	Q	L	Y	A	V	V	S	-	E	E	P	I	Y	hFgr	
287	F	L	E	E	A	N	L	M	K	T	L	Q	H	D	K	L	V	R	L	Y	A	V	V	T	R	E	E	P	I	Y	hLyn	
281	F	L	A	E	A	N	V	M	K	T	L	Q	H	D	K	L	V	K	L	H	A	V	V	T	K	E	-	P	I	Y	hHck	
285	F	L	A	E	A	N	L	M	K	Q	L	Q	H	Q	R	L	V	R	L	Y	A	V	V	T	-	Q	E	P	I	Y	hLck	
275	F	L	G	E	A	N	V	M	K	T	L	Q	H	E	R	L	V	R	L	Y	A	V	V	T	R	E	-	P	I	Y	mBik	

FIG.11C

304	I	I	T	E	L	M	R	H	G	S	L	Q	E	Y	L	Q	N	D	T	G	S	K	I	H	L	T	Q	Q	V	D	MKK3 MPI	aa
340	I	V	T	E	Y	M	N	K	G	S	L	L	D	F	L	K	D	G	E	G	R	A	L	K	L	P	N	L	V	D	hFyn	
339	I	V	T	E	F	M	S	Q	G	S	L	L	D	F	L	K	D	G	D	G	R	Y	L	K	L	P	Q	L	V	D	cYrk	
339	I	V	T	E	Y	M	S	K	G	S	L	L	D	F	L	K	G	E	T	G	K	Y	L	R	L	P	Q	L	V	D	hSrc	
346	I	V	T	E	F	M	S	K	G	S	L	L	D	F	L	K	E	G	D	G	K	Y	L	K	L	P	Q	L	V	D	hYes	
332	I	V	T	E	F	M	C	H	G	S	L	L	D	F	L	K	N	P	E	G	Q	D	L	R	L	P	Q	L	V	D	hFgr	
317	I	I	T	E	Y	M	A	K	G	S	L	L	D	F	L	K	S	D	E	G	G	K	V	L	L	P	K	L	I	D	hLyn	
310	I	I	T	E	F	M	A	K	G	S	L	L	D	F	L	K	S	D	E	G	S	K	Q	P	L	P	K	L	I	D	hHck	
314	I	I	T	E	Y	M	E	N	G	S	L	V	D	F	L	K	T	P	S	G	I	K	L	T	I	N	K	L	L	D	hLck	
304	I	V	T	E	Y	M	A	R	G	C	L	L	D	F	L	K	T	D	E	G	S	R	L	S	L	P	R	L	I	D	mBlk	

334	M	A	A	Q	V	A	S	G	M	A	Y	L	E	S	R	N	Y	I	H	R	D	L	A	A	R	N	V	L	V	G	MKK3 MPI	aa
370	M	A	A	Q	V	A	A	G	M	A	Y	I	E	R	M	N	U	I	H	R	D	L	R	S	A	N	I	L	V	G	hFyn	
369	M	A	A	Q	I	A	A	G	M	A	Y	I	E	R	M	N	Y	I	H	R	D	L	R	A	A	N	I	L	V	G	cYrk	
369	M	A	A	Q	I	A	S	G	M	A	Y	V	E	R	M	N	Y	V	H	R	D	L	R	A	A	N	I	L	V	G	hSrc	
376	M	A	A	Q	I	A	D	G	M	A	Y	I	E	R	M	N	Y	I	H	R	D	L	R	A	A	N	I	L	V	G	hYes	
362	M	A	A	Q	V	A	E	G	M	A	Y	M	E	R	M	N	Y	I	H	R	D	L	R	A	A	N	I	L	V	G	hFgr	
347	F	S	A	Q	I	A	E	G	M	A	Y	I	E	R	K	N	Y	I	H	R	D	L	R	A	A	N	V	L	V	S	hLyn	
340	F	S	A	Q	I	A	E	G	M	A	F	I	E	Q	R	N	Y	I	H	R	D	L	R	A	A	N	I	L	V	S	hHck	
344	M	A	A	Q	I	A	E	G	M	A	F	I	E	E	R	N	Y	I	H	R	D	L	R	A	A	N	I	L	V	S	hLck	
334	M	S	A	Q	V	A	E	G	M	A	Y	I	E	R	M	N	S	I	H	R	D	L	R	A	A	N	I	L	V	S	mBlk	

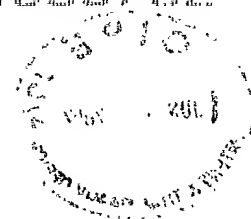
364	E	H	N	I	Y	K	V	A	D	F	G	L	A	R	V	F	K	V	D	N	E	D	I	Y	E	S	R	H	E	I	MKK3 MPI	aa
400	N	G	L	I	C	K	I	A	D	F	G	L	A	R	L	I	-	-	-	E	D	N	E	Y	T	A	R	Q	G	A	hFyn	
399	D	N	L	V	C	K	I	A	D	F	G	L	A	R	L	I	-	-	-	E	D	N	E	Y	T	A	R	Q	G	A	cYrk	
399	E	N	L	V	C	K	V	A	D	F	G	L	A	R	L	I	-	-	-	E	D	N	E	Y	T	A	R	Q	G	A	hSrc	
406	E	N	L	V	C	K	I	A	D	F	G	L	A	R	L	I	-	-	-	E	D	N	E	Y	T	A	R	Q	G	A	hYes	
392	E	R	L	A	C	K	I	A	D	F	G	L	A	R	L	I	-	-	-	K	D	E	Y	N	P	C	Q	G	S	hFgr		
377	E	S	L	M	C	K	I	A	D	F	G	L	A	R	V	I	-	-	-	E	D	N	E	Y	T	A	R	E	G	A	hLyn	
370	A	S	L	V	C	K	I	A	D	F	G	L	A	R	V	I	-	-	-	E	D	N	E	Y	T	A	R	E	G	A	hHck	
374	D	T	L	S	C	K	I	A	D	F	G	L	A	R	L	I	-	-	-	E	D	N	E	Y	T	A	R	E	G	A	hLck	
364	E	T	L	C	C	K	I	A	D	F	G	L	A	R	I	I	-	-	-	-	D	S	E	Y	T	A	Q	E	G	A	mBlk	

394	K	L	P	V	K	W	T	A	P	E	A	I	R	S	N	K	F	S	I	K	S	D	V	W	S	F	G	I	L	L	MKK3 MPI	aa
427	K	F	P	I	K	W	T	A	P	E	A	A	L	Y	G	R	F	T	I	K	S	D	V	W	S	F	G	I	L	L	hFyn	
426	K	F	P	I	K	W	T	A	P	E	A	A	L	F	G	K	F	T	I	K	S	D	V	W	S	F	G	I	L	L	cYrk	
426	K	F	P	I	K	W	T	A	P	E	A	A	L	Y	G	R	F	T	I	K	S	D	V	W	S	F	G	I	L	L	hSrc	
433	K	F	P	I	K	W	T	A	P	E	A	A	L	Y	G	R	F	T	I	K	S	D	V	W	S	F	G	I	L	Q	hYes	
419	K	F	P	I	K	W	T	A	P	E	A	A	L	F	G	R	F	T	I	K	S	D	V	W	S	F	G	I	L	L	hFgr	
404	K	F	P	I	K	W	T	A	P	E	A	I	N	F	G	C	F	T	I	K	S	D	V	W	S	F	G	I	L	L	hLyn	
397	K	F	P	I	K	W	T	A	P	E	A	I	N	F	G	S	F	T	I	K	S	D	V	W	S	F	G	I	L	L	hHck	
401	K	F	P	I	K	W	T	A	P	E	A	I	N	Y	G	T	F	T	I	K	S	D	V	W	S	F	G	I	L	L	hLck	
390	K	F	P	I	K	W	T	A	P	E	A	I	H	F	G	V	F	T	I	K	A	D	V	W	S	F	G	V	L	L	mBlk	

FIG.11D

Title: NOVEL
MEGAKARYOCYTIC PROTEIN
TYROSINE KINASES
Inventor(s): Axel ULLRICH et al.
Appl. No.: 09/977,261

09/977261 060702



424	Y	E	I	I	T	Y	G	K	M	P	Y	S	G	M	T	G	A	Q	V	I	Q	M	L	A	Q	N	Y	R	L	P	MKK3 MPI aa
457	T	E	L	V	T	K	G	R	V	P	Y	P	G	M	N	N	R	E	V	L	E	Q	V	E	R	G	Y	R	M	P	hFyn
456	T	E	L	V	T	K	G	R	V	P	Y	P	G	M	N	N	R	E	V	L	E	Q	V	E	R	G	Y	R	M	Q	cYrk
456	T	E	L	T	T	K	G	R	V	P	Y	P	G	M	V	N	R	E	V	L	D	Q	V	E	R	G	Y	R	M	P	hSrc
463	T	E	L	V	T	K	G	R	V	P	Y	P	G	M	V	N	R	E	V	L	E	Q	V	E	R	G	Y	R	M	P	hYes
449	T	E	L	I	T	K	G	R	I	P	Y	P	G	M	N	K	R	E	V	L	E	Q	V	E	Q	G	Y	H	M	P	hFgr
434	Y	E	I	V	T	Y	G	K	I	P	Y	P	G	R	T	N	A	D	V	M	T	A	L	S	Q	G	Y	R	M	P	hLyn
427	M	E	I	V	T	Y	G	R	I	P	Y	P	G	M	S	N	P	E	V	I	R	A	L	E	R	G	Y	R	M	P	hHck
431	T	E	I	V	T	H	G	R	I	P	Y	P	G	M	T	N	P	E	V	I	Q	N	L	E	R	G	Y	R	M	V	hLck
420	M	V	I	V	T	Y	G	R	V	P	Y	P	G	M	S	N	P	E	V	I	R	S	L	E	H	G	Y	R	M	P	mB1k
454	Q	P	S	N	C	P	Q	Q	F	Y	N	-	I	M	L	E	C	W	N	A	E	P	K	E	R	P	T	F	E	T	MKK3 MPI aa
487	C	P	Q	D	C	P	I	S	L	H	-	E	L	M	I	H	C	W	K	K	D	P	E	E	R	P	T	F	E	Y	hFyn
486	C	P	G	G	C	P	P	S	L	H	-	D	V	M	V	Q	C	W	K	R	E	P	E	E	R	P	T	F	E	Y	cYrk
486	C	P	P	E	C	P	E	S	L	H	-	D	L	M	C	Q	C	W	R	K	E	P	E	E	R	P	T	F	E	Y	hSrc
493	C	P	Q	G	C	P	E	S	L	H	-	E	L	M	N	L	C	W	K	K	D	P	D	E	R	P	T	F	E	Y	hYes
479	C	P	P	G	C	P	A	S	L	Y	-	E	A	M	E	Q	T	W	R	L	D	P	E	E	R	P	T	F	E	Y	hFgr
464	R	V	E	N	C	P	D	E	L	Y	-	D	I	M	K	M	C	W	K	E	K	A	E	E	R	P	T	F	D	Y	hLyn
457	R	P	E	N	C	P	E	E	L	Y	-	N	I	M	M	R	C	W	K	N	R	P	E	E	R	P	T	F	E	Y	hHck
461	R	P	D	N	C	P	E	E	L	Y	-	Q	L	M	R	L	C	W	K	E	R	P	E	D	R	P	T	F	D	Y	hLck
450	C	P	E	T	C	P	P	E	L	Y	N	D	I	I	T	E	C	W	R	G	R	P	E	E	R	P	T	F	E	F	mB1k
483	L	R	W	K	L	E	D	Y	F	E	-	T	D	S	S	Y	S	D	A	N	N	F	I	R							MKK3 MPI aa
516	L	Q	S	F	L	E	D	Y	F	T	A	T	E	P	Q	Y	Q	P	G	E	N	-	-	-	L						hFyn
515	L	Q	S	F	L	E	D	Y	F	T	A	T	E	P	Q	Y	Q	P	G	D	N	-	-	-	Q						cYrk
515	L	Q	A	F	L	E	D	Y	F	T	S	T	E	P	Q	Y	Q	P	G	E	N	-	-	-	L						hSrc
522	I	Q	S	F	L	E	D	Y	F	T	A	T	E	P	Q	Y	Q	P	G	E	N	-	-	-	L						hYes
508	L	Q	S	F	L	E	D	Y	F	T	S	A	E	P	Q	Y	Q	P	G	D	Q	-	-	-	T						hFgr
493	L	Q	S	V	L	D	D	F	Y	T	A	T	E	G	Q	Y	Q	Q	-	-	Q	-	-	-	P						hLyn
486	I	Q	S	V	L	D	D	F	Y	T	A	T	E	S	Q	Y	Q	Q	-	-	Q	-	-	-	P						hHck
490	L	R	S	V	L	E	D	F	F	T	A	T	E	G	Q	Y	Q	P	-	-	Q	-	-	-	P						hLck
480	L	Q	S	V	L	E	D	F	Y	T	A	T	E	G	Q	Y	E	L	-	-	Q	-	-	-	P						mB1k

FIG.11E

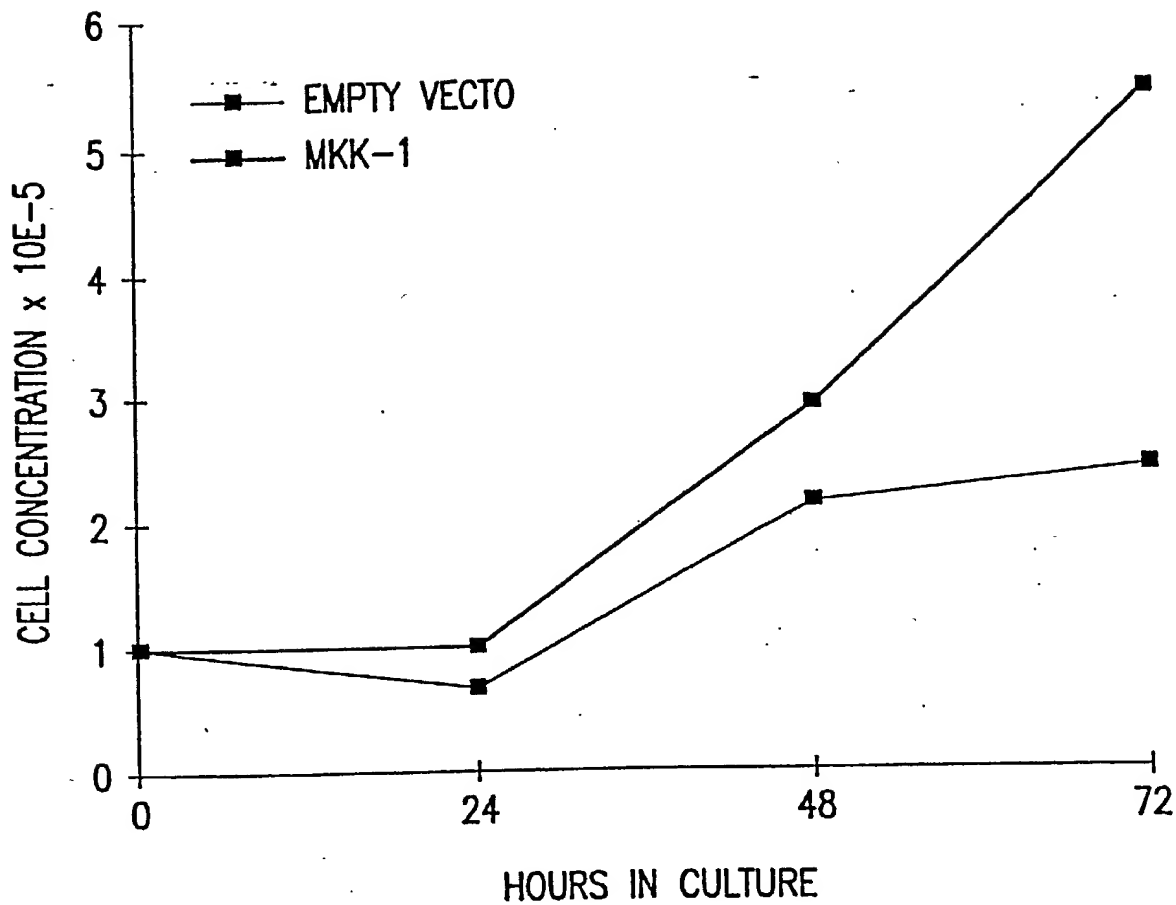


FIG.12

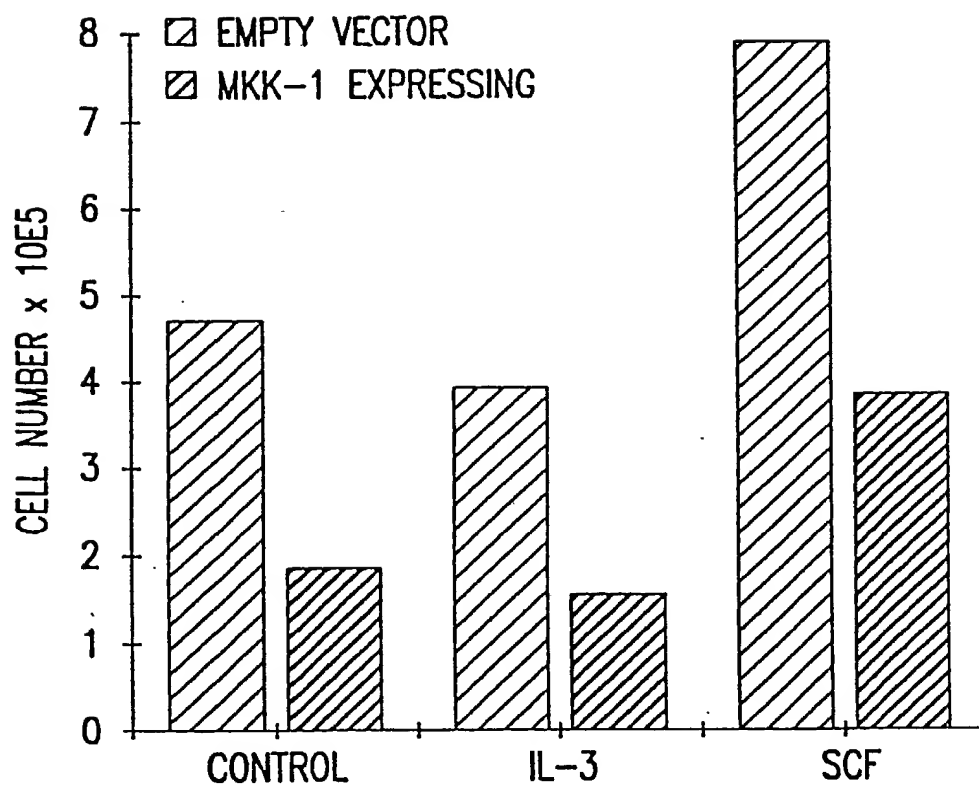
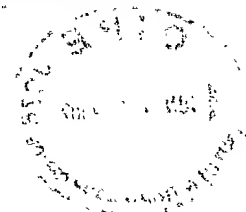


FIG.13

Title: NOVEL
MEGAKARYOCYTIC PROTEIN
TYROSINE KINASES
Inventor(s): Axel ULLRICH et al.
Appl. No.: 09/977,261

09/977,261 .060702

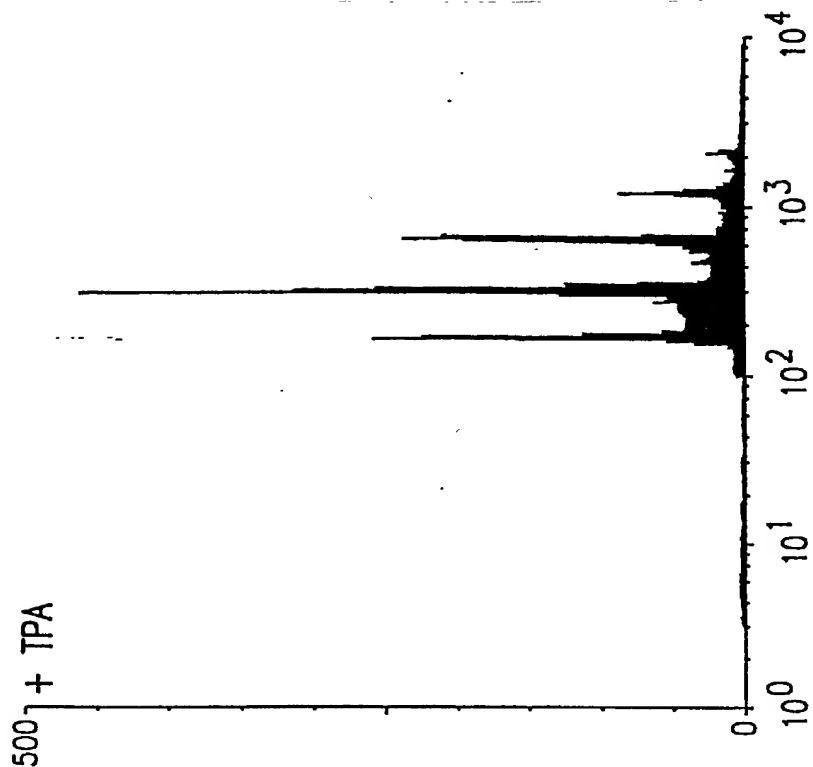
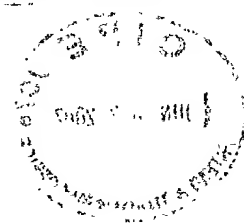


FIG.14B

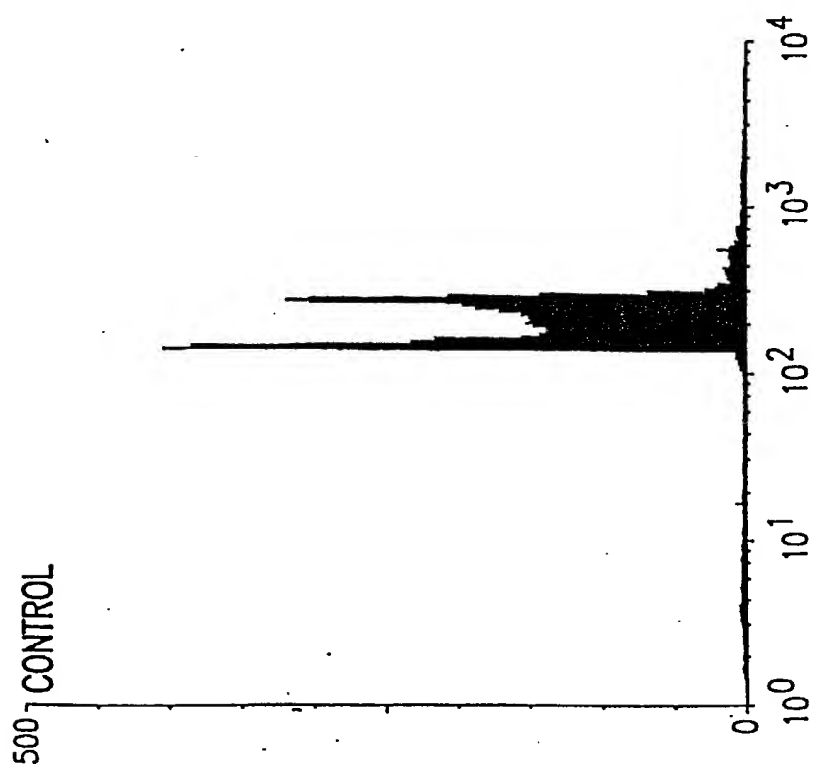


FIG.14A

Title: NOVEL
MEGAKARYOCYTIC PROTEIN
TYROSINE KINASES
Inventor(s): Axel ULLRICH et al.
Appl. No.: 09/977,261

09/977,261 .060702

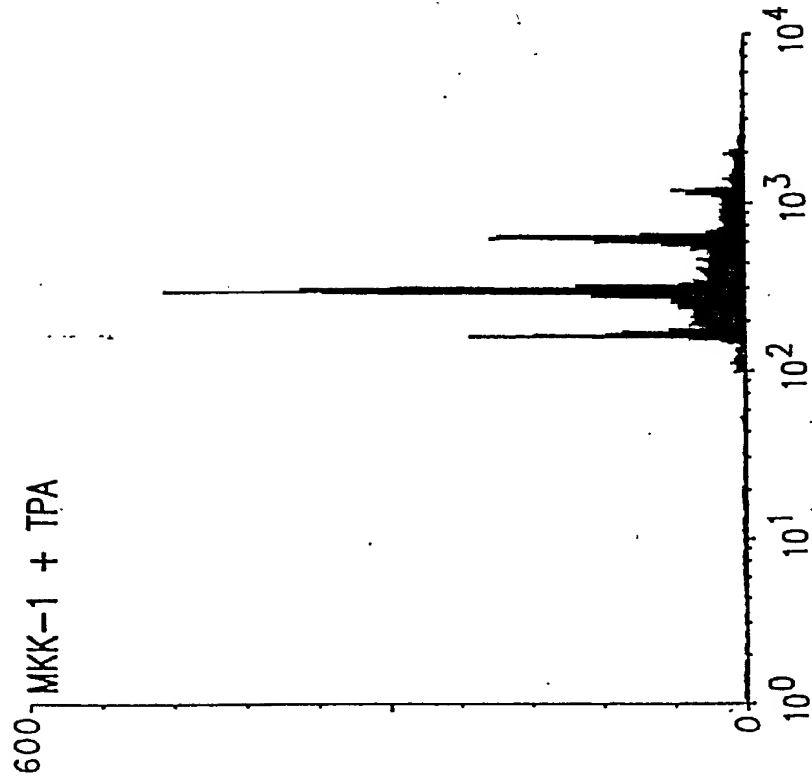


FIG.14D

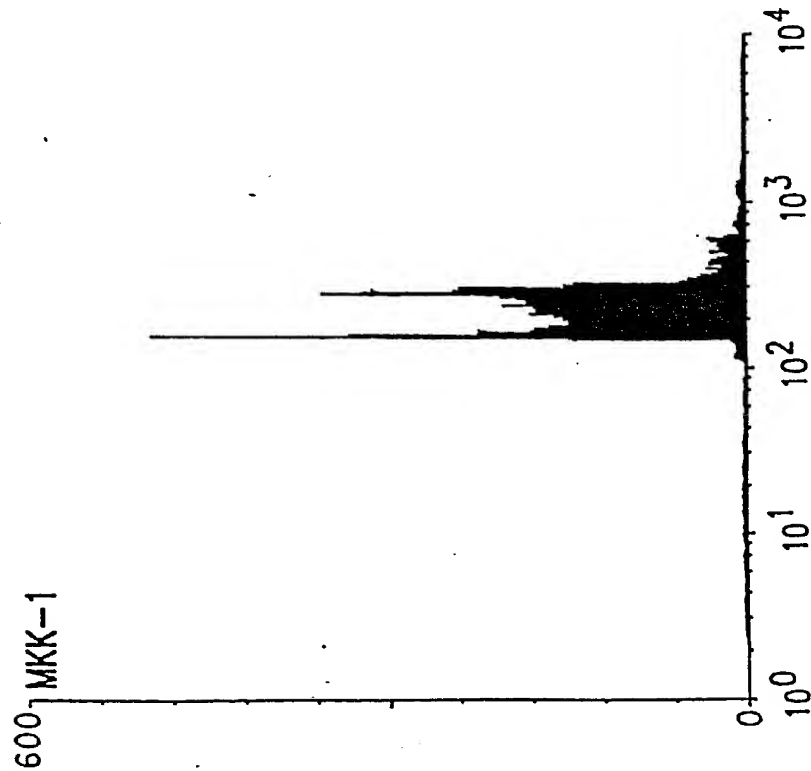


FIG.14C